


```
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541 aagagaatca tacagtggct ttattcttac tacttaaaaa aaggatgatgt gatggcagtg
601 atggtaaca tcacacaggg aagacc
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//

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CGCTCAGGATAC GACTTCGTCGCTAGAGGATCGGATCCCGGCGGATTATATAGCTCGATCGATCT
 TTCTCTATAT CCGCGGATATGGGATATATACACACACACCGCGGATAGCATGACTGATCTA
 ECGCATCTATCT
 CACAGACTCTACCGCT

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Structure

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Details

Display **GenBank** Show:

1: AI692778. wd70d04.x1 NCI_CG...[gi:4970118]

Links

LOCUS AI692778 565 bp mRNA linear EST 17-DEC-1999
 DEFINITION wd70d04.x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:2336935 3',
 mRNA sequence.

ACCESSION AI692778

VERSION AI692778.1 GI:4970118

KEYWORDS EST.

SOURCE Homo sapiens (human)

ORGANISM Homo sapiens

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
 Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 565)

AUTHORS NCI-CGAP <http://www.ncbi.nlm.nih.gov/ncicgap>.

TITLE National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
 Tumor Gene Index

JOURNAL Unpublished (1997)

COMMENT Contact: Robert Strausberg, Ph.D.

Email: cgapbs-r@mail.nih.gov

Tissue Procurement: Christopher Moskaluk, M.D., Ph.D., Michael R.
 Emmert-Buck, M.D., Ph.D.

cDNA Library Preparation: M. Bento Soares, Ph.D.

cDNA Library Arrayed by: Greg Lennon, Ph.D.

DNA Sequencing by: Washington University Genome Sequencing Center

Clone distribution: NCI-CGAP clone distribution information can be
 found through the I.M.A.G.E. Consortium/LLNL at:

www-bio.llnl.gov/bbrp/image/image.html

Insert Length: 637 Std Error: 0.00

Seq primer: -40UP from Gibco

High quality sequence stop: 487.

FEATURES Location/Qualifiers

source

1..565

/organism="Homo sapiens"

/db_xref="taxon:9606"

/clone="IMAGE:2336935"

/clone_lib="NCI_CGAP_Lu24"

/tissue_type="carcinoid"

/lab_host="DH10B"

/note="Organ: lung; Vector: pT7T3D-Pac (Pharmacia) with a
 modified polylinker; Plasmid DNA from the normalized
 library NCI_CGAP_Lu5 was prepared, and ss circles were
 made in vitro. Following HAP purification, this DNA was
 used as tracer in a subtractive hybridization reaction.
 The driver was PCR-amplified cDNAs from a pool of 5,000
 clones made from the same library (cloneIDs
 1414920-1417991 and 1520904-1522439). Subtraction by Bento
 Soares and M. Fatima Bonaldo."

BASE COUNT 142 a 146 c 168 g 109 t

ORIGIN

1 gcggccgcag cgctgaatgg ggtggaccga cgttcctcgc agcggtcagc aaggctggct
 61 ctagaagtgc tggagagggc caagaggagg gcggtggact ggcattgcct ggagcgctcc
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541 tgaccatcac tgccatcaca tcacc
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